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1600

RAW SEQUENCE LISTING

DATE: 01/28/2003

PATENT APPLICATION: US/09/529,239D

TIME: 08:42:38

Input Set : A:\09529239ThirdSubSeqList.txt

Output Set: N:\CRF4\01282003\I529239D.raw

1 <110> APPLICANT: Doutriaux, Marie-Pascale
 2 Betzner, Andreas
 3 Freyssinet, Georges
 4 Perez, Pascal
 6 <120> TITLE OF INVENTION: METHOD FOR OBTAINING PLANT VARIETIES
 9 <130> FILE REFERENCE: A33153-PCT-USA 072667.0128
 11 <140> CURRENT APPLICATION NUMBER: 09/529,239D
 12 <141> CURRENT FILING DATE: 2000-10-27
 14 <150> PRIOR APPLICATION NUMBER: PCT/EP98/06977
 15 <151> PRIOR FILING DATE: 1998-10-09
 17 <160> NUMBER OF SEQ ID NOS: 100
 19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
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 23 <212> TYPE: DNA
 24 <213> ORGANISM: Artificial Sequence
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 27 <221> NAME/KEY: modified_base
 28 <222> LOCATION: (11)...(11) ✓
 29 <223> OTHER INFORMATION: I
 31 <221> NAME/KEY: modified_base
 32 <222> LOCATION: (14)...(14)
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 36 <222> LOCATION: (17)...(17)
 37 <223> OTHER INFORMATION: I
 39 <223> OTHER INFORMATION: Degenerate oligonucleotides UPMU used to isolate
 40 AtMSH3 and AtMSH6.
 42 <300> PUBLICATION INFORMATION:
 43 <301> AUTHORS: Reenan and Kolodner
 44 <302> TITLE: Genetics
 45 <303> JOURNAL: 132
 46 <306> PAGES: 963-973
 47 <307> DATE: 1992
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 W--> 50 ctggatccac nggncnaay atg 23
 52 <210> SEQ ID NO: 2
 53 <211> LENGTH: 23
 54 <212> TYPE: DNA
 55 <213> ORGANISM: Artificial Sequence
 57 <220> FEATURE:
 58 <223> OTHER INFORMATION: Degenerate oligonucleotides DOMU used to isolate
 59 AtMSH3 and AtMSH6.

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      63 <223> OTHER INFORMATION: I
W--> 65 <221> modified_base
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      67 <223> OTHER INFORMATION: I
      69 <300> PUBLICATION INFORMATION:
      70 <301> AUTHORS: Reenan and Kolodner
      71 <302> TITLE: Genetics
      72 <303> JOURNAL: 132
      73 <306> PAGES: 963-973
      74 <307> DATE: 1992
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W--> 77 ctggatccrt artgngtnrc raa 23
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      82 <213> ORGANISM: Artificial Sequence
      84 <220> FEATURE:
      85 <223> OTHER INFORMATION: MSH3 specific primer 636 for PCR using cDNA of
      86 Arabidopsis thaliana ecotype Columbia.
      89 <400> SEQUENCE: 3
      90 tgctagtgcc tcttgcaagc tcat 24
      92 <210> SEQ ID NO: 4
      93 <211> LENGTH: 27
      94 <212> TYPE: DNA
      95 <213> ORGANISM: Artificial Sequence
      97 <220> FEATURE:
      98 <223> OTHER INFORMATION: Primer AP1 for PCR using cDNA of Arabidopsis
      99 thaliana ecotype Columbia containing adapter
     100 sequences ligated to both its ends.
     102 <400> SEQUENCE: 4
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     107 <212> TYPE: DNA
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     124 <223> OTHER INFORMATION: MSH3 specific primer S525 for PCR using cDNA of

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132 <212> TYPE: DNA
133 <213> ORGANISM: Artificial Sequence
135 <220> FEATURE:
136 <223> OTHER INFORMATION: MSH3 specific primer S51 for PCR using cDNA of
137      Arabidopsis thaliana ecotype Columbia.
139 <400> SEQUENCE: 7
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142 <210> SEQ ID NO: 8
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148 <223> OTHER INFORMATION: MSH3 specific primer 635 for PCR using cDNA of
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159 <220> FEATURE:
160 <223> OTHER INFORMATION: MSH3 specific primer S523 for PCR using cDNA of
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168 <212> TYPE: DNA
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171 <220> FEATURE:
172 <223> OTHER INFORMATION: MSH3 specific primer 1S5 for PCR using cDNA of
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183 <220> FEATURE:
184 <223> OTHER INFORMATION: MSH3 specific primer S53 for PCR using cDNA of
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188 gacaaagagc gaaatgaggc ccotttg          27
190 <210> SEQ ID NO: 12

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202 tccccgactc acgaaccgaa tccggtagcc gaatcatcaa caccgccacc gaagatatcc 120
203 gccactgtat ccttctctcc ttccaagcgt aagcttctct ccgaccacct cgccgccgcg 180
204 tcacccaaaa agcctaaact ttctcctcac actcaaaacc cagtaccoga tcccaattta 240
205 caccaaagat ttctccagag atttctggaa ccctcgccgg aggaatatgt tcccgaacg 300
206 tcatcatcga ggaaatacac accattggaa cagcaagtgg tggagctaaa gagcaagtac 360
207 ccagatgtgg tttgatggg ggaagtggg tacaggtaca gattcttcgg agaagacgcg 420
208 gagatcgtag cacgcgtgtt gggatattac gtcatatgg atcacaattt catgacggcg 480
209 agtgtgccaa catttcgatt gaatttccat gtgagaagac tggatgaatgc aggatacaag 540
210 attggtgtag tgaagcagac tgaaactgca gccattaagt cccatggtgc aaaccggacc 600
211 ggcccttttt tccggggact gtcggcggtg tataccaaaag ccacgcttga agcggctgag 660
212 gatataagtg gtggttgtgg tggatgaaga ggttttgggt cacagagtaa tttcttgggt 720
213 tgtgttgtgg atgagagagt taagtgcggg acattaggct gtggtattga aatgagtttt 780
214 gatgttagag tcggtgttgt tggcggtgaa atttcgacag gtgaagttgt ttatgaagag 840
215 ttcaatgata atttcatgag aagtggatta gaggtgtga ttttgagctt gtcaccagct 900
216 gagctgttgc ttggccagcc tctttcaca caaactgaga agtttttggg ggcacatgct 960
217 ggacctacct caaacgttcg agtggaaact gcctcactgg attgtttcag caatggtaat 1020
218 gcagtagatg aggttatttc attatgtgaa aaaatcagcg caggtaactt agaagatgat 1080
219 aaagaaatga agctggaggc tgctgaaaaa ggaatgtctt gcttgacagt tcatacaatt 1140
220 atgaacatgc cacatctgac tgttcaagcc ctgcacctaa cgttttgcca tctcaaacag 1200
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225 <212> TYPE: DNA
226 <213> ORGANISM: Artificial Sequence
228 <220> FEATURE:
229 <223> OTHER INFORMATION: MSH3 specific primer 2S5 for PCR using cDNA of
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232 <400> SEQUENCE: 13
233 atccccgggtc aaaatgaaca agttggtttt agtc 34
235 <210> SEQ ID NO: 14
236 <211> LENGTH: 27
237 <212> TYPE: DNA
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: MSH3 specific primer S52 for PCR using cDNA of
242 Arabidopsis thaliana ecotype Columbia.
244 <400> SEQUENCE: 14
245 gccacatctg actgttcaag ccctcgc 27
247 <210> SEQ ID NO: 15
248 <211> LENGTH: 2110

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Input Set : A:\09529239ThirdSubSeqList.txt

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259 tgaaaggatc ctttaccaag gggcctcatt tcgctctttg tcaagtaaca cagagatgac 120
260 tctctcagcc aatactctgc aacagtttga ggttggtgaa aataattcag atggatcgga 180
261 atctggctcc ttattccata atatgaatca cacacttaca gtatatggtt ccaggcttct 240
262 tagacactgg gtgactcatc ctctatgcga tagaaatttg atatctgctc ggcttgatgc 300
263 tgtttctgag atttctgctt gcatgggata tcatagttct tcccagctca gcagtgaagt 360
264 ggttgaagaa ggttctgaga gagcaattgt atcacctgag ttttatctcg tgctctcttc 420
265 agtcttgaca gctatgtcta gatcatctga tattcaacgt ggaataacaa gaatctttca 480
266 tcggactgct aaagccacag agttcattgc agttatggaa gctattttac ttgcggggaa 540
267 gcaaattcag cggcttggca taaagcaaga ctctgaaatg aggagtatgc aatctgcaac 600
268 tgtgcgatct actcttttga gaaaattgat ttctgttatt tcatcccctg ttgtggttga 660
269 caatgccgga aaacttctct ctgccctaaa taaggaagcg gctgttcgag gtgacttgct 720
270 cgacatacta atcacttcca gcgaccaatt tcctgagctt gctgaagctc gccaaagcag 780
271 tttagtcatc agggaaaagc tggattcctc gatagcttca tttcgcaaga agctcgctat 840
272 tcgaaatttg gaatttcttc aagtgtcggg gatcacacat ttgatagagc tgcccgttga 900
273 ttccaaggtc cctatgaatt gggtgaaagt aaatagcacc aagaagacta ttcgatatca 960
274 tccccagaa atagtagctg gcttgatga gctagctcta gcaactgaac atcttgccat 1020
275 tgtgaaccga gcttcgtggg atagtttctt caagagtttc agtagatact acacagattt 1080
276 taaggctgcc gttcaagctc ttgctgcaat ggactgtttg cactcccttt caactctatc 1140
277 tagaaacaag aactatgtcc gtcccagagt tgtggatgac tgtgaaccag ttgagataaa 1200
278 catacagtct ggtcgtcatc ctgtactgga gactatatta caagataact tcgtcccaaa 1260
279 tgacacaatt ttgcatgcag aaggggaata ttgccaaatt atcaccggac ctaacatggg 1320
280 aggaaagagc tgctatatcc gtcaagttgc tttaatttcc ataattggctc aggttggttc 1380
281 ctttgtacca cgtctattcg ccaagctgca cgtgcttgat ggtgttttca ctcggtggg 1440
282 tgcttcagac agtatccagc atggcagaag tacctttcta gaagaattaa gtgaagcgtc 1500
283 acacataatc agaacctgtt cttctcgttc gcttggtata ttagatgagc ttggaagagg 1560
284 cactagcaca cagcaggtg tagccattgc ctatgcaaca ttacagcatc tcctagcaga 1620
285 aaagagatgt ttggttcttt ttgtcacgca ttaccctgaa atagctgaga tcagtaacgg 1680
286 attcccaggt tctgttgga cataccatgt ctcgatctg acattgcaga aggataaagg 1740
287 cagttatgat catgatgat tgacctacct atataagctt gtgcgtggtc tttgcagcag 1800
288 gagctttggt ttttaagggt ctcagcttgc ccagatacct ccatcatgta tacgtcgagc 1860
289 catttcaatg gctgcaaaat tggaaagctga ggtacgtgca agagagagaa atacacgcat 1920
290 gggagaacca gaaggacatg aagaaccgag aggcgcagaa gaatctatct cggctctagg 1980
291 tgacttggtt gcagacctga aatttgctct ctctgaagag gaccttgga aagcattcga 2040
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293 ttgacccggg                                     2110
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296 <211> LENGTH: 29
297 <212> TYPE: DNA
298 <213> ORGANISM: Artificial Sequence
300 <220> FEATURE:
301 <223> OTHER INFORMATION: MSH3 specific primer S51 for PCR using cDNA of

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 11, 14, 17
Seq#:2; N Pos. 15, 18